

CORRIGENDUM

## Whole-exome sequencing in 20,197 persons for rare variants in Alzheimer's disease

In Raghavan et al. (2018)<sup>1</sup>, Endnote citations from a previous draft of the paper were incorporated into the variant numbers of three of the variants, making these variant numbers incorrect. The corrected Table 3 is provided below. We apologize for this error.

**Table 3.** SORL1 variants.

Genomic Position	Variant Type	Variant Class	CADD score	Protein modification	ExAC Global Frequency	Case/Control	Sex	Ethnicity	Braak Stage	Age at Onset or Last Visit
11-121367577	snv	SAV	26.6	NA	0	case	F	AA	NA	77
11-121367654	snv	SG	37	p.Arg279*	0	case	F	NHW	6	72
<b>11-121421343<sup>23</sup></b>	snv	SG	39	p.Arg744*	0	case	M	NHW	NA	65
<b>11-121421343<sup>23</sup></b>	snv	SG	39	p.Arg744*	0	case	F	NHW	NA	67
11-121426001	indel	FV	NA	p.Asp850fs	0	case	F	NHW	NA	60
11-121428047	snv	SG	41	p.Arg866*	0	case	M	NHW	6	65
11-121430263	indel	FV	NA	p.Ile983fs	0	ctrl	M	AA	NA	64
11-121440980	snv	SDV	27.6	NA	4.95E-05	case	F	CH	NA	80
11-121456930	snv	SAV	26.8	NA	0	case	M	NHW	NA	69
11-121456930	snv	SAV	26.8	NA	0	case	M	NHW	6	62
11-121461788	indel	FV	NA	p.Cys1431fs	0	case	F	NHW	NA	61
<b>11-121466482<sup>25</sup></b>	snv	SDV	28	NA	0	case	F	NHW	3	90+
<b>11-121466482<sup>25</sup></b>	snv	SDV	28	NA	0	case	F	NHW	NA	90+
11-121474911	indel	FV	NA	p.Thr1511fs	0	case	M	NHW	NA	60
11-121474984	snv	SG	35	p.Cys1534*	0	case	F	NHW	NA	74
<b>11-121477568<sup>25</sup></b>	snv	SG	46	p.Arg1655*	0	case	M	NHW	NA	69
11-121477667	snv	SDV	26.9	NA	0	case	F	AA	NA	68
11-121485637	indel	FV	NA	p.Asp1828fs	0	case	M	NHW	NA	75
11-121491801	indel	FV	NA	p.Lys1975fs	0	case	M	NHW	6	61
11-121500253	indel	FV	NA	p.Met2211fs	0	case	M	NHW	6	62

Those in bold have previously been identified as indicated by the reference

SNV, Single Nucleotide Variant; Indel, Insertion or Deletion; CADD, Combined Annotation Dependent Depletion; FV, Frameshift Variant; SAV, Splice Acceptor Variant; SDV, Splice Donor Variant; SG, Stop Gained; AA, African American; CH, Caribbean Hispanic; NHW, Non-hispanic White

### Reference

- Raghavan NS, Brickman AM, Andrews H, Manly JJ, Schupf N, Lantigua R, Wolock CJ, Kamalakaran S, Petrovski S, Tosto G, Vardarajan BN, Goldstein DB, Mayeux R and Alzheimer's Disease Sequencing Project. Whole-exome sequencing in 20,197 persons for rare variants in Alzheimer's disease. *Ann Clin Transl Neurol* 2018;5:832–842. <https://doi.org/10.1002/acn3.582>.